

Accepted

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RAW SEQUENCE LISTING
PATENT APPLICATION US/09/491,577

 DATE: 02/15/2000
 TIME: 14:09:10

Input Set: I491577.RAW

This Raw Listing contains the General Information Section and up to first 5 pages.

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1 <110> APPLICANT: Yale University
2           Carlson, John R.
3           Kim, Hunhyong
4           Clyne, Peter J.
5           Warr, Coral G.
6 <120> TITLE OF INVENTION: Novel Family of Odorant Receptor Genes in Drosophila
7 <130> FILE REFERENCE: 44574-5061-US
8 <140> CURRENT APPLICATION NUMBER: US/09/491,577
9 <141> CURRENT FILING DATE: 2000-01-25
10 <150> EARLIER APPLICATION NUMBER: US 60/117,132
11 <151> EARLIER FILING DATE: 1999-01-25
12 <160> NUMBER OF SEQ ID NOS: 112
13 <170> SOFTWARE: PatentIn Ver. 2.1
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27      cggtt aag tcc cga gat gcc ttc gtt tac tta gat cggtt atg tgg    96
28      Arg Val Lys Ser Arg Asp Ala Phe Val Tyr Leu Asp Arg Val Met Trp
29          20          25          30
30      tcc ttt ggc tgg aca gtg cct gaa aac aaa agg tgg gat cta cat tac   144
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32          35          40          45
33      aaa ctg tgg tca act ttc gtg aca ttg ttg ata ttt atc ctt ctg ccg   192
34      Lys Leu Trp Ser Thr Phe Val Thr Leu Leu Ile Phe Ile Leu Leu Pro
35          50          55          60
36      ata tcg gta agc gtt gag tat att cag cgg ttc aag acc ttc tcg gcg   240
37      Ile Ser Val Ser Val Glu Tyr Ile Gln Arg Phe Lys Thr Phe Ser Ala
38          65          70          75          80
39      ggt gag ttt ctt agc tca atc cag att ggc gtt aac atg tac gga agc   288
40      Gly Glu Phe Leu Ser Ser Ile Gln Ile Gly Val Asn Met Tyr Gly Ser
41          85          90          95
42      agc ttt aaa agt tat ttg acc atg atg gga tat aag aag aga cag gag   336
43      Ser Phe Lys Ser Tyr Leu Thr Met Met Gly Tyr Lys Lys Arg Gln Glu
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63	ttg atc tcc atg gta ata gca cga tgc cac atc acc ctt ctg aaa cag	672
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 99 Val Asp Pro Glu Lys Gln Phe Tyr Ile Ser Ser Ile Ala Glu Val Ile
 100 180 185 190
 101 Leu Arg Gly Trp Ala Val Phe Met Asp Leu Cys Thr Asp Val Cys Pro
 102 195 200 205
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 104 210 215 220
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 130 50 55 60
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 136 85 90 95
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 140 agc ctg ttg ttg ctc agc tct ggc acg gcg aca aat gcc gcc ttc acc 384
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 142 115 120 125
 143 ttg caa ccc ctg att atg ggt ctc tac cgc tgg att gtg cag ctg cca 432
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150	Gln Pro Gly Val Phe Pro Leu Thr Tyr Val Leu Leu Thr Ala Ser Gly			
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152	gcc tgc acc gtt ttc gcc ttc agc ttc gtg gac gga ttc ttc att tgc			576
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154	180 185 190			
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201 35 40 45
202 Phe Ala Met Val Val Val Gly Ala Val Gly Glu Val Ser Tyr Gly Cys
203 50 55 60
204 Val His Leu Asp Asn Leu Val Val Ala Leu Glu Ala Phe Cys Pro Gly
205 65 70 75 80
206 Thr Thr Lys Ala Val Cys Val Leu Lys Leu Trp Val Phe Phe Arg Ser
207 85 90 95
208 Asn Arg Arg Trp Ala Glu Leu Val Gln Arg Leu Arg Ala Ile Leu Leu
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210 Ser Leu Leu Leu Ser Ser Gly Thr Ala Thr Asn Ala Ala Phe Thr
211 115 120 125
212 Leu Gln Pro Leu Ile Met Gly Leu Tyr Arg Trp Ile Val Gln Leu Pro
213 130 135 140
214 Gly Gln Thr Glu Leu Pro Phe Asn Ile Ile Leu Pro Ser Phe Ala Val
215 145 150 155 160
216 Gln Pro Gly Val Phe Pro Leu Thr Tyr Val Leu Leu Thr Ala Ser Gly
217 165 170 175
218 Ala Cys Thr Val Phe Ala Phe Ser Phe Val Asp Gly Phe Phe Ile Cys
219 180 185 190
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**VERIFICATION SUMMARY
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